



FIG. 1

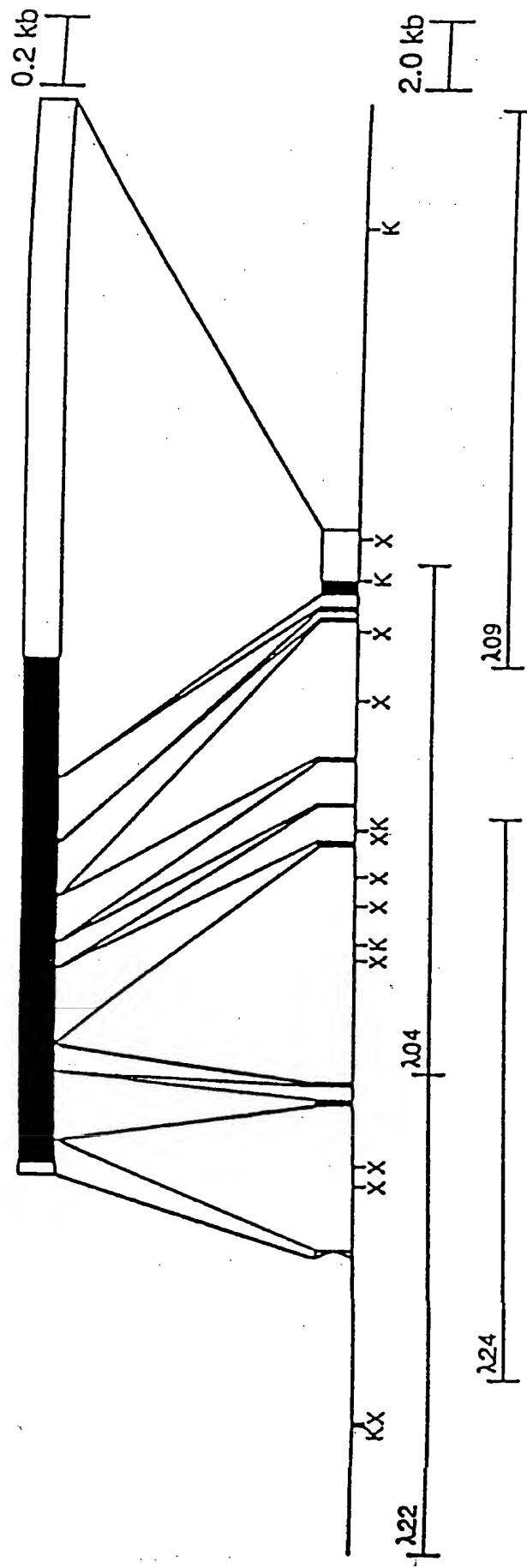


FIG. 2

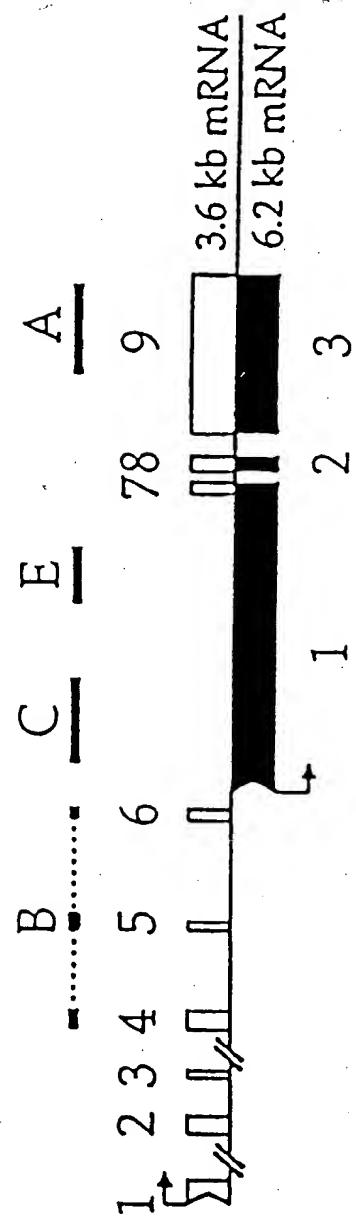


FIG. 3

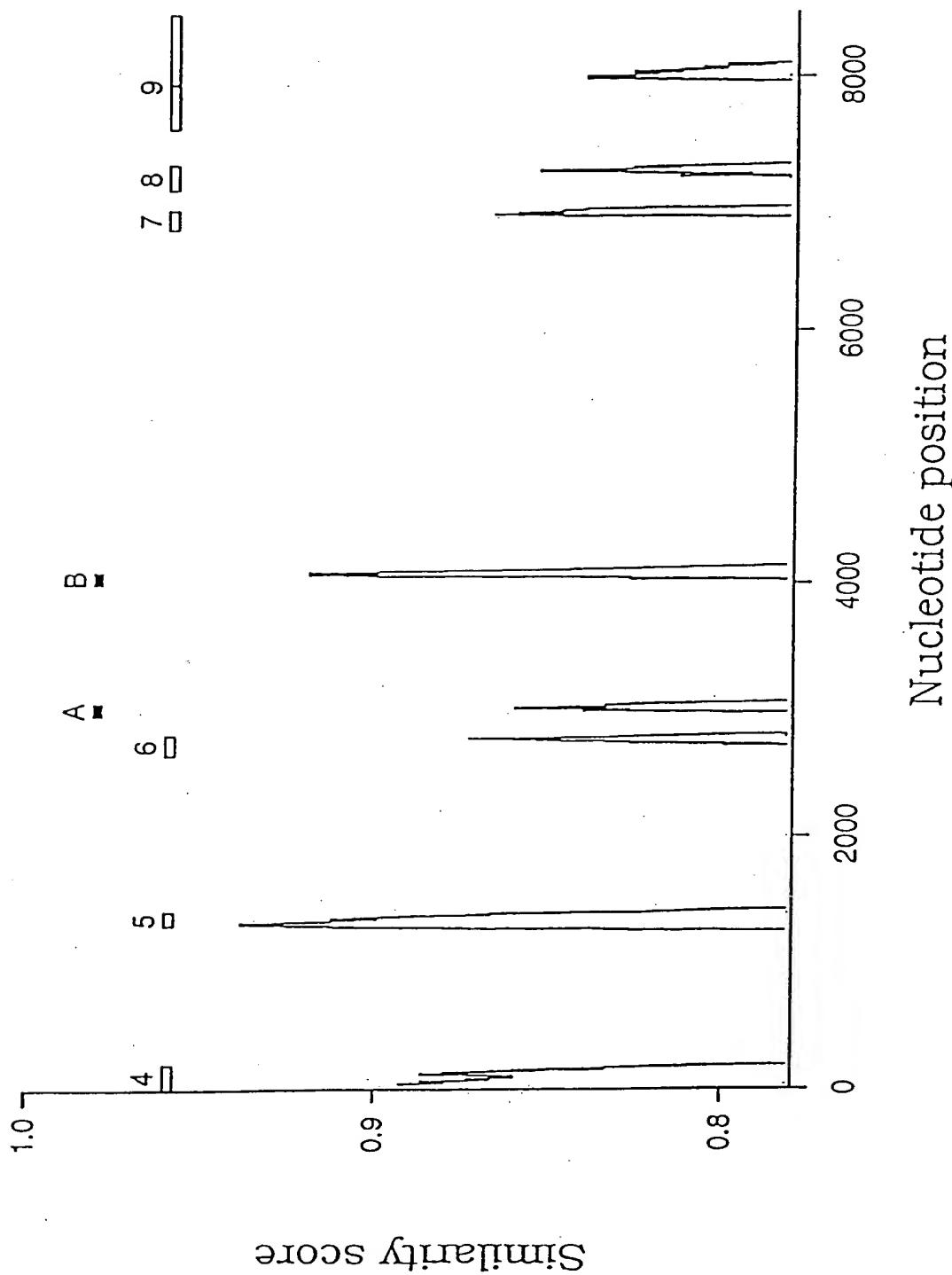
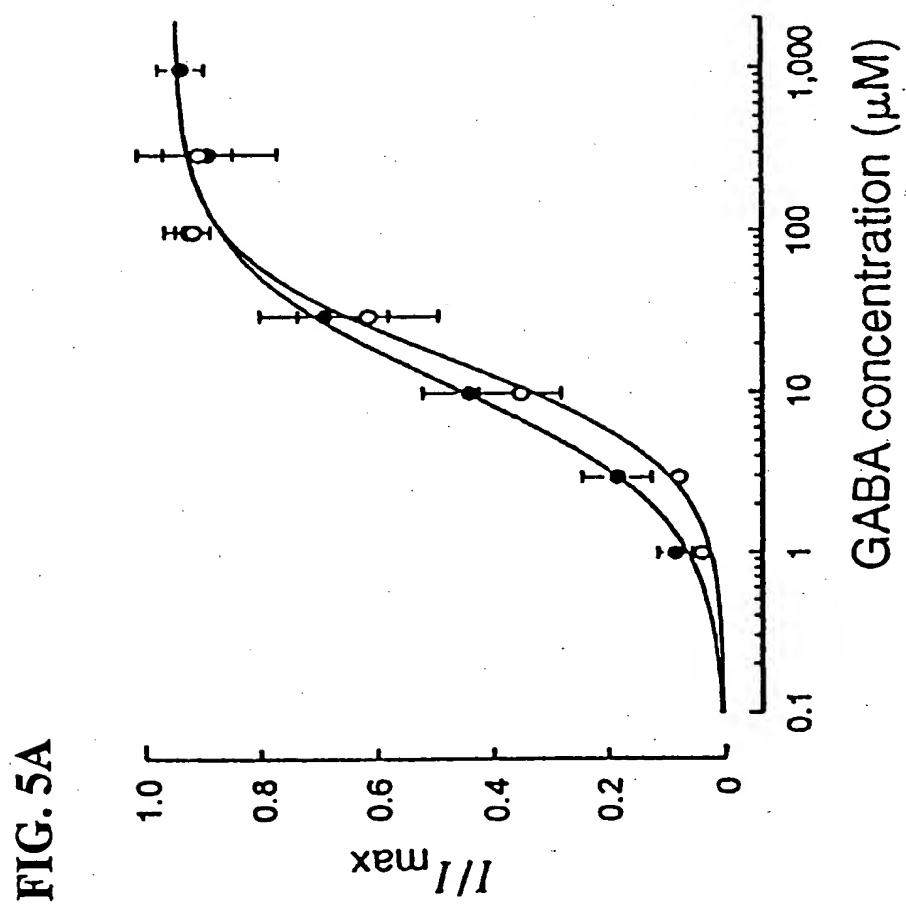


FIG. 4

| | | |
|-----------|---|-----|
| ε subunit | <u>MLSKVLFVLLGILLILQSRVEGPQTESKNEASSR</u> DVYGPQPQLENQLLSEETKSTETETGSRVGKLP EASRIL | 75 |
| ε subunit | <u>NTILSNYDHKLRLRGIGEKPTVVTVEIAVNSLGPLSIL</u> DMEYTIDIIIFSQTWYDERLCYNDT FESLVLNGNVSQL | 150 |
| Consensus | L Y RP G S M Y Q W D R L L | |
| ε subunit | <u>WIPDTFFRNSKRTHETHEITMPNQMVRIYKDGKVLYTIRMTIDAGC</u> SLHMLRFPMDSHSCPLSESSES SYPENEMTY | 225 |
| Consensus | W PDT N H T N R G LY R T C L P D C L SY Y | |
| ε subunit | <u>KWENFKLEINEKNISWKL</u> FQFDFTGVSNKTEIIITPVGDFMNTIFFNVSRREGYVAFQNYVPSST VTFMLSWVSEFW | 300 |
| Consensus | W L Q G F L R G Q Y P S VSEFW | |
| ε subunit | <u>IKTESAPARTSLGITSVLTMTLGTFSRKNFPRVSYITALDFYIAICF</u> VFCALLEFAVLFNLI YNOTKAHASP | 375 |
| Consensus | AR G TIVLMTT R LP A D C FVFALE | |
| ε subunit | <u>KLRHPRINSRAHARTRARSACARQHQEAFTVCQIVTTEGSDGERPSCSAQQPPSPGSPEGPRSLCSKLIACCEWC</u> | 450 |
| Consensus | | |
| ε subunit | <u>KRFKKYFCMVPDCEGSTWQQGRRLCIHVYRLDNYSRVVFPTTEFFENVLYMLVCLNL</u> | 506 |
| Consensus | D R FP F N YW Y | |



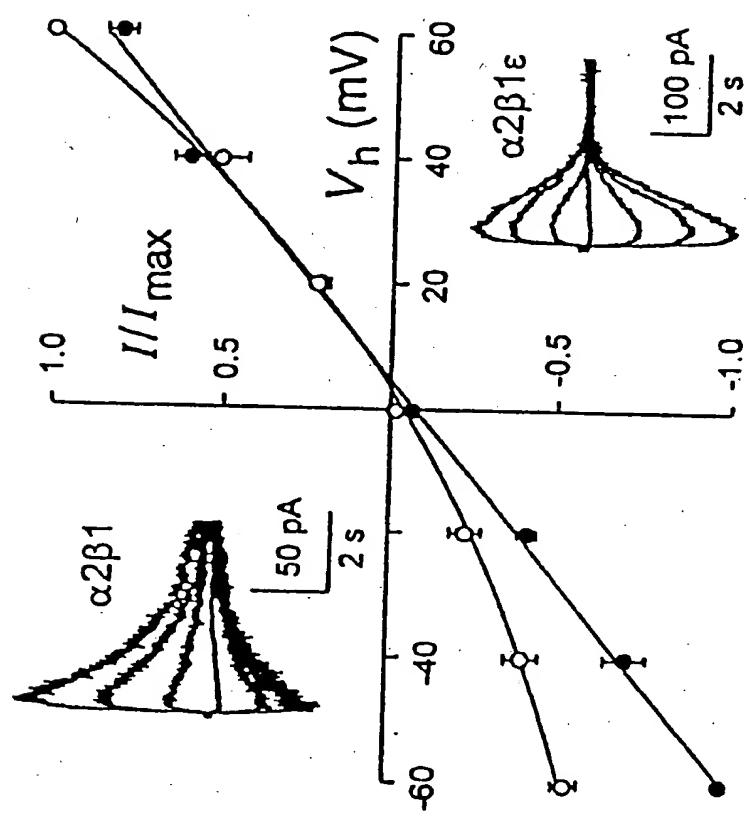


FIG. 5B

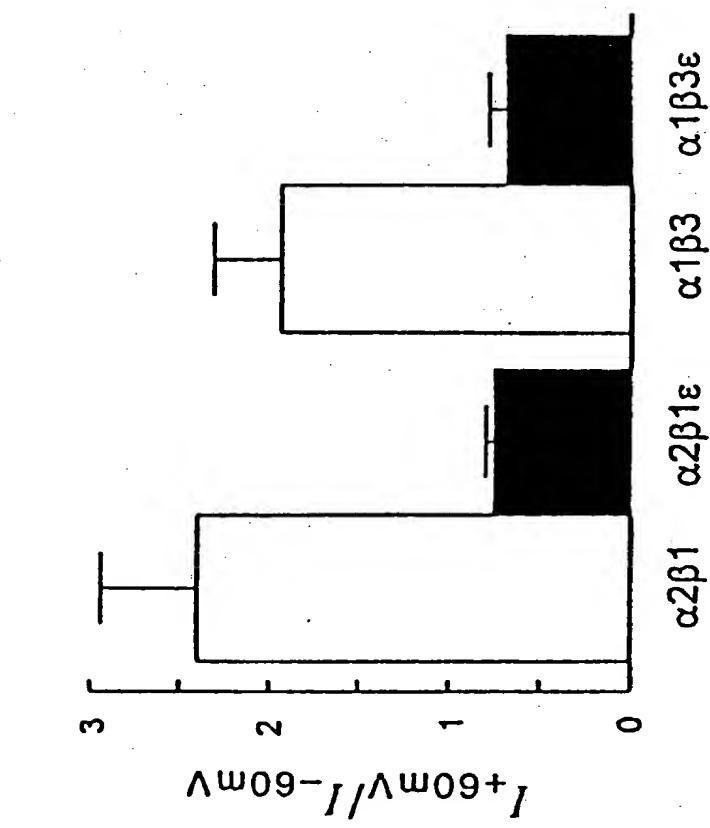


FIG. 5C

FIG. 6A

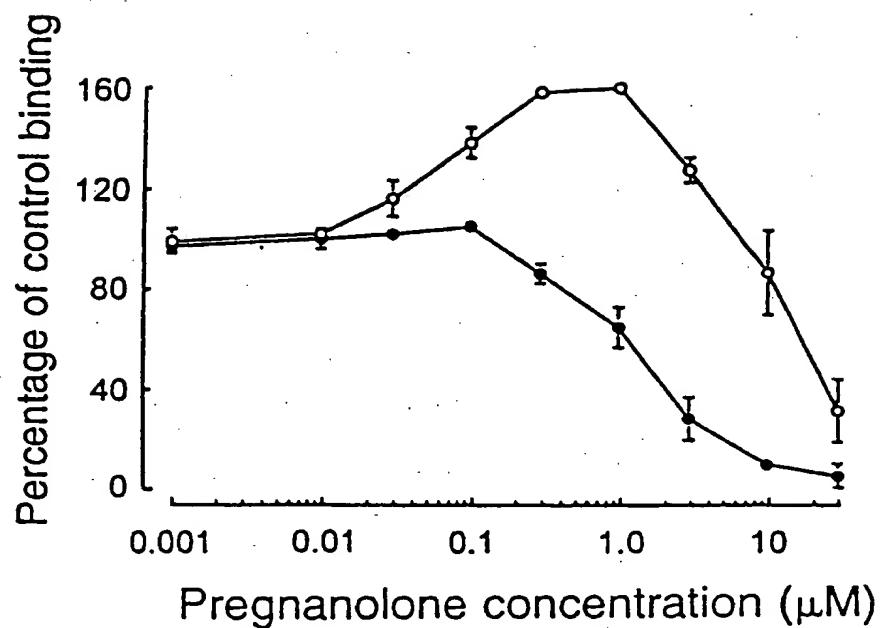


FIG. 6B

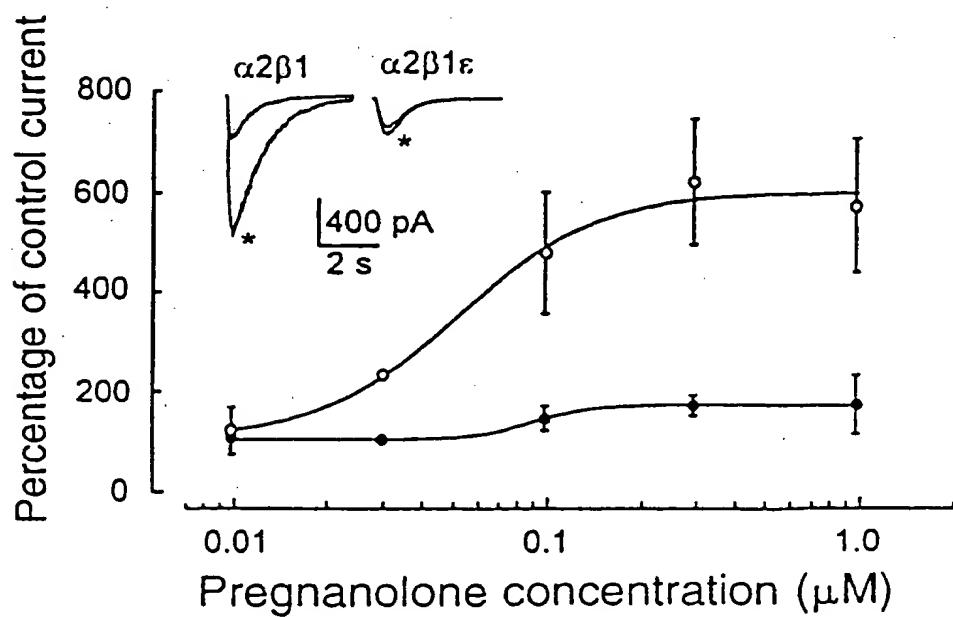


FIG. 6C

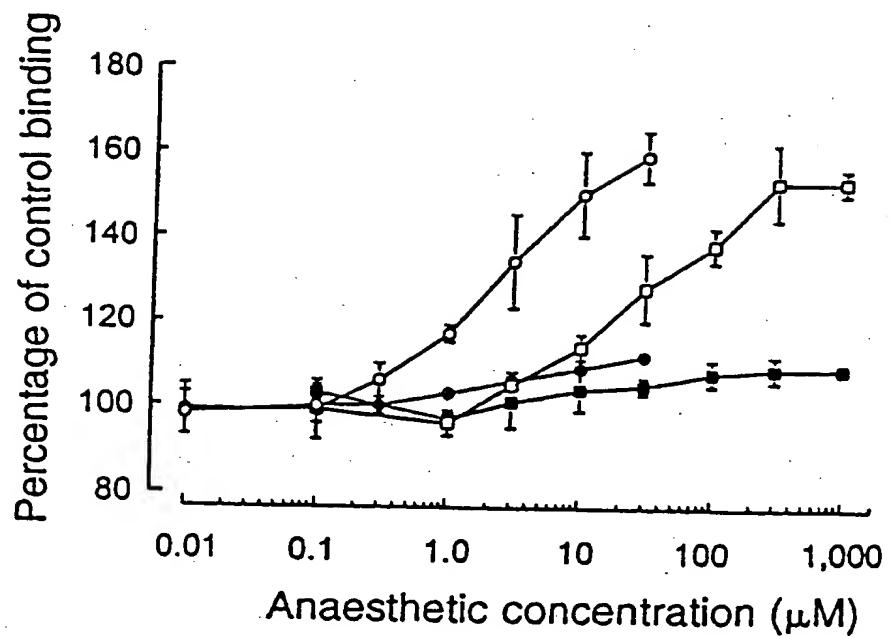
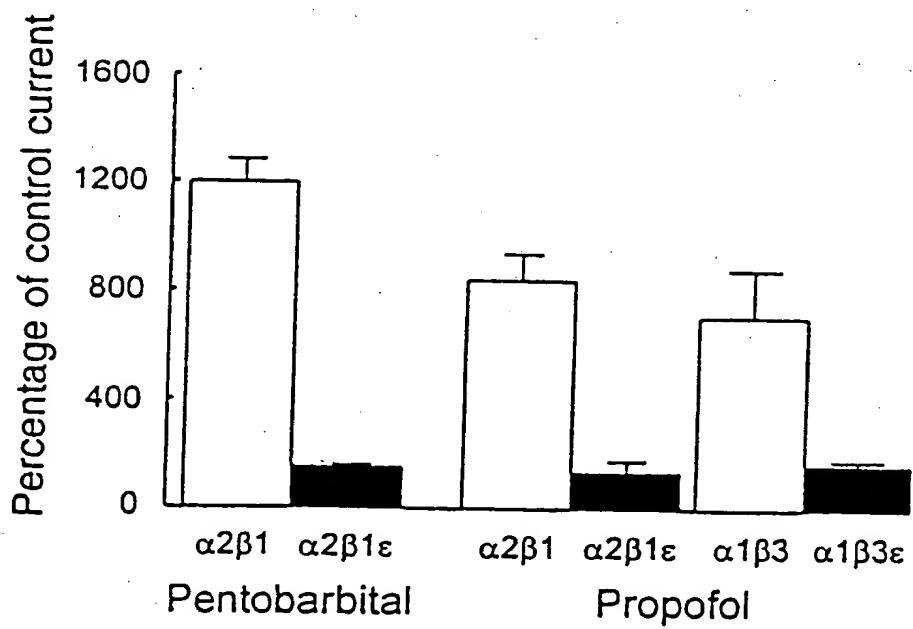


FIG. 6D



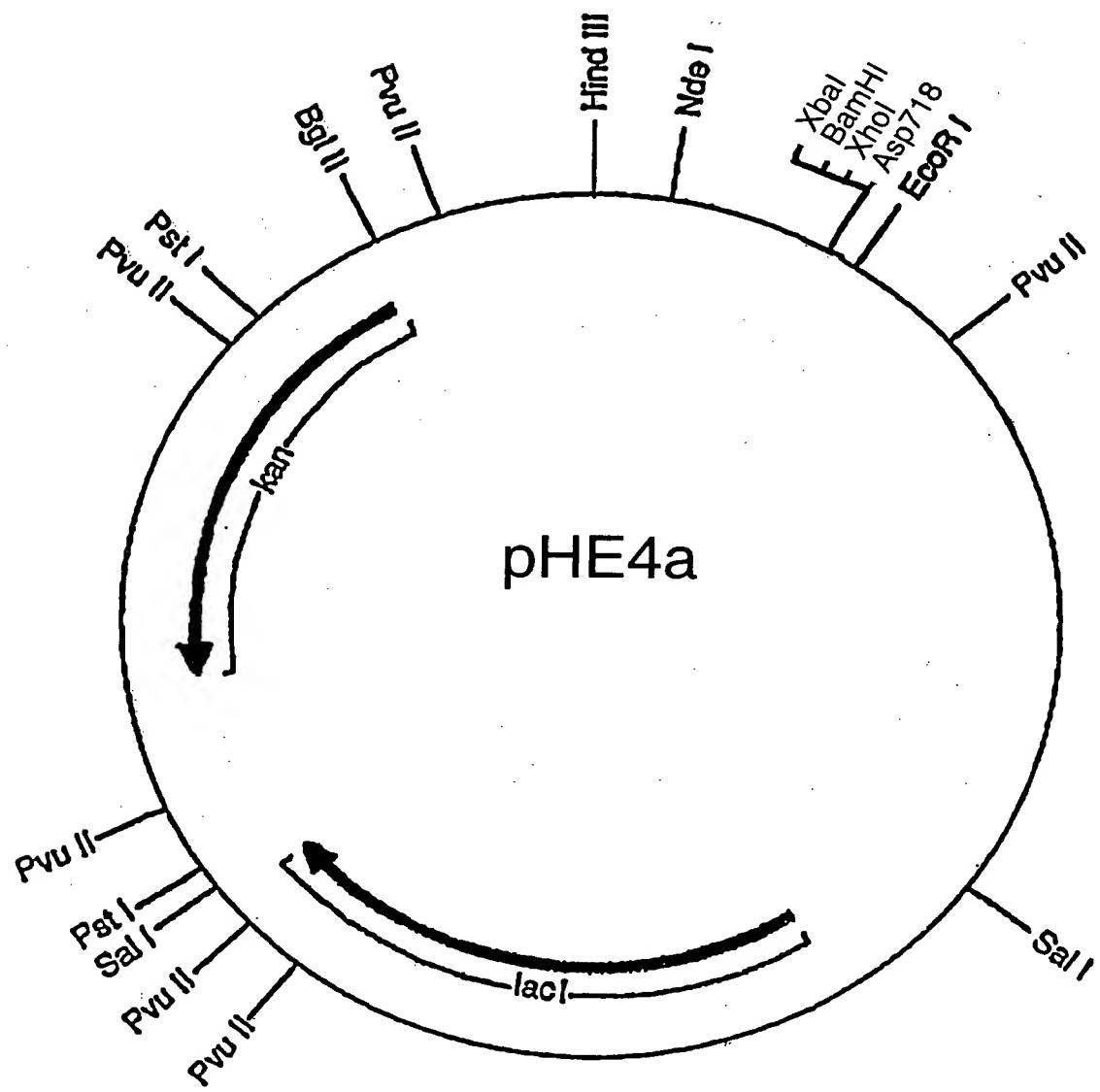


FIG. 7

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The diagram illustrates the SV40 early promoter region with transcription start sites and operators. The sequence is oriented 5' to 3' from top to bottom.

- Operator 1:** Located at position -35, consisting of the sequence **TITGACT** (boxed).
- Operator 2:** Located at position -10, consisting of the sequence **TTGAGGAAACATAACAAAT** (boxed).
- S/D:** Located at position 50, consisting of the sequence **TAAGATGTACCCAATTGAGGGAAACAAAT** (boxed).
- Transcription Start:** Indicated by a vertical arrow at position 94, pointing to the sequence **AGAGGAGAAATCATATG** (boxed).

Below the sequence, the positions are labeled: -35, Operator 1, -10, Operator 2, 50, S/D, and 94.